

OIPF

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/842,469

DATE: 05/11/2001
 TIME: 11:51:23

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3 <110> APPLICANT: PFIZER PRODUCT INC
 5 <120> TITLE OF INVENTION: ADAMTS POLYPEPTIDES, NUCLEIC ACIDS ENCODING THEM, AND
 6 USES THEREOF
 8 <130> FILE REFERENCE: PC10473B
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/842,469
 C--> 11 <141> CURRENT FILING DATE: 2001-04-26
 13 <160> NUMBER OF SEQ ID NOS 9
 15 <170> SOFTWARE: PatentIn Ver. 2.1
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 3776
 19 <212> TYPE: DNA
 20 <213> ORGANISM: nucleotide human
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 26 cgtgcacacg tggcctctat ggtccgca tgcagatcc tcggtgggc cctggccctg 180
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59 ggcgaatgca agcactggg ctgcgacga gtcttggtgt ccgacctgcg ggagacaaag 2220
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88 <210> SEQ ID NO: 2

89 <211> LENGTH: 1104

90 <212> TYPE: PRT

91 <213> ORGANISM: amino acid/human

92 <400> SEQUENCE: 2

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95 Gly Leu Met Phe Glu Val Thr His Ala Phe Arg Ser Gln Asp Glu Phe

96 20 25 30

97 Leu Ser Ser Leu Gln Ser Tyr Glu Ile Ala Phe Pro Thr Arg Val Asp

98 35 40 45

99 His Asn Gly Ala Leu Leu Ala Phe Ser Pro Pro Pro Pro Arg Arg Gln

100 50 55 60

101 Arg Arg Gly Thr Gly Ala Ile Ala His Ser Arg Leu Phe Tyr Cys Val

102 65 70 75 80

103 Ala Ser Pro Ser Thr His Phe Leu Leu Asn Leu Thr Arg Ser Ser Arg

104 85 90 95

105 Leu Leu Ala Gly His Val Ser Val Glu Tyr Trp Thr Arg Glu Gly Leu

106 100 105 110

107 Ala Trp Gln Arg Ala Ala Arg Pro His Cys Leu Tyr Ala Gly His Leu

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118 Gln Gly Gln Ala Ser Thr Ser His Val Ala Ile Ser Thr Cys Gly Gly
119          130          135          140
121 Leu His Gly Leu Ile Val Ala Asp Glu Glu Glu Tyr Leu Ile Glu Pro
122 145          150          155          160
124 Leu His Gly Gly Pro Lys Gly Ser Arg Ser Pro Glu Glu Ser Gly Pro
125          165          170          175
127 His Val Val Tyr Lys Arg Ser Ser Leu Arg His Pro His Leu Asp Thr
128          180          185          190
130 Ala Cys Gly Val Arg Asp Glu Lys Pro Trp Lys Gly Arg Pro Trp Trp
131          195          200          205
133 Leu Arg Thr Leu Lys Pro Pro Pro Ala Arg Pro Leu Gly Asn Glu Thr
134          210          215          220
136 Glu Arg Gly Gln Pro Gly Leu Lys Arg Ser Val Ser Arg Glu Arg Tyr
137 225          230          235          240
139 Val Glu Thr Leu Val Val Ala Asp Lys Met Met Val Ala Tyr His Gly
140          245          250          255
142 Arg Arg Asp Val Glu Gln Tyr Val Leu Ala Ile Met Asn Ile Val Ala
143          260          265          270
145 Lys Leu Phe Gln Asp Ser Ser Leu Gly Ser Thr Val Asn Ile Leu Val
146          275          280          285
148 Thr Arg Leu Ile Leu Leu Thr Glu Asp Gln Pro Thr Leu Glu Ile Thr
149          290          295          300
151 His His Ala Gly Lys Ser Leu Asp Ser Phe Cys Lys Trp Gln Lys Ser
152 305          310          315          320
154 Ile Val Asn His Ser Gly His Gly Asn Ala Ile Pro Glu Asn Gly Val
155          325          330          335
157 Ala Asn His Asp Thr Ala Val Leu Ile Thr Arg Tyr Asp Ile Cys Ile
158          340          345          350
160 Tyr Lys Asn Lys Pro Cys Gly Thr Leu Gly Leu Ala Pro Val Gly Gly
161          355          360          365
163 Met Cys Glu Arg Glu Arg Ser Cys Ser Val Asn Glu Asp Ile Gly Leu
164          370          375          380
166 Ala Thr Ala Phe Thr Ile Ala His Glu Ile Gly His Thr Phe Gly Met
167 385          390          395          400
169 Asn His Asp Gly Val Gly Asn Ser Cys Gly Ala Arg Gly Gln Asp Pro
170          405          410          415
172 Ala Lys Leu Met Ala Ala His Ile Thr Met Lys Thr Asn Pro Phe Val
173          420          425          430
175 Trp Ser Ser Cys Ser Arg Asp Tyr Ile Thr Ser Phe Leu Asp Ser Gly
176          435          440          445
178 Ser Gly Leu Cys Leu Asn Asn Arg Phe Ile Ala Gln Asp Phe Val Tyr
179          450          455          460
181 Pro Thr Val Ala Pro Gly Gln Ala Tyr Asn Ala Asp Ile Gln Cys Arg
182 465          470          475          480
184 Phe Gln His Gly Val Lys Ser Arg Gln Cys Lys Tyr Gly Glu Val Cys
185          485          490          495
187 Ser Glu Leu Trp Cys Leu Ser Lys Ser Asn Arg Cys Ile Thr Asn Ser
188          500          505          510

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191      515      520      525
193 Gly Trp Cys Tyr Lys Arg Val Cys Val Pro Phe Gly Ser Arg Pro Glu
194      530      535      540
196 Gly Val Asp Gly Ala Trp Gly Pro Trp Thr Pro Trp Gly Asp Cys Ser
197 545      550      555      560
199 Arg Thr Cys Gly Gly Val Ser Ser Ser Ser Arg His Cys Asp Ser
200      565      570      575
202 Pro Arg Pro Thr Ile Gly Gly Lys Tyr Cys Leu Gly Glu Arg Arg Arg
203      580      585      590
205 His Arg Ser Cys Asn Thr Asp Asp Cys Pro Pro Gly Ser Gln Asp Phe
206      595      600      605
208 Arg Glu Val Gln Cys Ser Glu Phe Asp Ser Ile Pro Phe Arg Gly Lys
209      610      615      620
211 Phe Tyr Lys Trp Lys Thr Tyr Arg Gly Gly Gly Val Lys Ala Cys Ser
212 625      630      635      640
214 Leu Thr Cys Leu Ala Glu Gly Phe Asn Phe Tyr Thr Glu Arg Ala Ala
215      645      650      655
217 Ala Val Val Asp Gly Thr Pro Cys Arg Pro Asp Thr Val Asp Ile Cys
218      660      665      670
220 Val Ser Gly Glu Cys Lys His Val Gly Cys Asp Arg Val Leu Gly Ser
221      675      680      685
223 Asp Leu Arg Glu Asp Lys Cys Arg Val Cys Gly Gly Asp Gly Ser Ala
224      690      695      700
226 Cys Glu Thr Ile Glu Gly Val Phe Ser Pro Ala Ser Pro Gly Ala Gly
227 705      710      715      720
229 Tyr Glu Asp Val Val Trp Ile Pro Lys Gly Ser Val His Ile Phe Ile
230      725      730      735
232 Gln Asp Leu Asn Leu Ser Leu Ser His Leu Ala Leu Lys Gly Asp Gln
233      740      745      750
235 Glu Ser Leu Leu Leu Glu Gly Leu Pro Gly Thr Pro Gln Pro His Arg
236      755      760      765
238 Leu Pro Leu Ala Gly Thr Thr Phe Gln Leu Arg Gln Gly Pro Asp Gln
239      770      775      780
241 Val Gln Ser Leu Glu Ala Leu Gly Pro Ile Asn Ala Ser Leu Ile Val
242 785      790      795      800
244 Met Val Leu Ala Arg Thr Glu Leu Pro Ala Leu Arg Tyr Arg Phe Asn
245      805      810      815
247 Ala Pro Ile Ala Arg Asp Ser Leu Pro Pro Tyr Ser Trp His Tyr Ala
248      820      825      830
250 Pro Trp Thr Lys Cys Ser Ala Gln Cys Ala Gly Gly Ser Gln Val Gln
251      835      840      845
253 Ala Val Glu Cys Arg Asn Glu Leu Asp Ser Ser Ala Val Ala Pro His
254      850      855      860
256 Tyr Cys Ser Ala His Ser Lys Leu Pro Lys Arg Gln Arg Ala Cys Asn
257 865      870      875      880
259 Thr Glu Pro Cys Pro Pro Asp Trp Val Val Gly Asn Trp Ser Leu Cys
260      885      890      895
262 Ser Arg Ser Cys Asp Ala Gly Val Arg Ser Arg Ser Val Val Cys Gln

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265 Arg Arg Val Ser Ala Ala Glu Glu Lys Ala Leu Asp Asp Ser Ala Cys
266          915          920          925
268 Pro Gln Pro Arg Pro Pro Val Leu Glu Ala Cys His Gly Pro Thr Cys
269          930          935          940
271 Pro Pro Glu Trp Ala Ala Leu Asp Trp Ser Glu Cys Thr Pro Ser Cys
272 945          950          955          960
274 Gly Pro Gly Leu Arg His Arg Val Val Leu Cys Lys Ser Ala Asp His
275          965          970          975
277 Arg Ala Thr Leu Pro Pro Ala His Cys Ser Pro Ala Ala Lys Pro Pro
278          980          985          990
280 Ala Thr Met Arg Cys Asn Leu Arg Arg Cys Pro Pro Ala Arg Trp Val
281          995          1000          1005
283 Ala Gly Glu Trp Gly Glu Cys Ser Ala Gln Cys Gly Val Gly Gln Arg
284          1010          1015          1020
286 Gln Arg Ser Val Arg Cys Thr Ser His Thr Gly Gln Ala Ser His Glu
287 1025          1030          1035          1040
289 Cys Thr Glu Ala Leu Arg Pro Pro Thr Thr Thr Gln Gln Lys Glu Ala
290          1045          1050          1055
292 Lys Cys Asp Ser Pro Thr Pro Gly Asp Gly Pro Glu Glu Cys Lys Asp
293          1060          1065          1070
295 Val Asn Cys Val Ala Tyr Cys Pro Leu Val Leu Lys Phe Gln Phe Cys
296          1075          1080          1085
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299          1090          1095          1100
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306 (211) LENGTH: 3414
307 (212) TYPE: DNA
308 (213) ORGANISM: nucleotide,mouse
310 (400) SEQUENCE: 3
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VERIFICATION SUMMARY

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L:11 M:271 C: Current Filing Date differs. Replaced Current Filing Date